

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

Notch(C)	IDEC-SNP	CONGCTC---D-VGSY-C-CPPGFT	GK---GE-N
10244(C)	-NECTM---	COH---C	VNT-GSY-CKC-SG--
42			G--L-C D
57	VNECGMKPRP	COHR C	CRCPGYT
108	VNSRTCAMIN	COYS C	EDTBEGPOCLCPSS
166	IDECASGKVI	CPYNRRC	VNTFGSYCKCHIGFE
212	INECTMDSHT	CSHANC	FNTQGSF CKCKQGYK
			GNGRLCS
CD97(C)	V-EC-SG-Q--C-SS--C	-NTVGSY-CRCRPGW-P-PG-PN---	D
EGF(C)	NSDSECP LSHDGYCLHDGVCMYIEALDKYACNCVGYI---GER--COYRDLKMWELR		

FIG. 1

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GGCTGGAGAA GAAACAGCAA GGGAGTCTGT GAAGCTACAT GCGAACCTGG  
ATGTAAGTTT GGTGAGTGCG TGGGACCAA CAAATGCAGA TGCTTTCCAG  
GATACACCGG GAAAACCTGC AGTCAAGATG TGAATGAGTG TGGAATGAAA  
CCCCGGCCAT GCCAACACAG ATGTGTGAAT ACACACGGAA GCTACAAGTG  
CTTTTGCCTC AGTGGCCACA TGCTCATGCC AGATGCTACG TGTGTGAACT  
CNAGGACATG TGCCATGATA AACTGTCAGT ATAGCTGTGA AGACACAGAA  
(SEQ ID NO 1)

GGCTGGAGAA GAAACAGCAA GGGAGTCTGT GAAGCTACAT GCGAACCTGG  
ATGTAAGTTT GGTGAGTGCG TGGGACCAA CAAATGCAGA TGCTTTCCAG  
GATACACCGG GAAAACCTGC AGTCAAGATG TGAATGAGTG TGGAATGAAA  
CCCCGGCCAT GCCAACACAG ATGTGTGAAT ACACACGGAA GCTACAAGTG  
CTTTTGCCTC AGTGGCCACA TGCTCATGCC AGATGCTACG TGTGTGAACT  
CNAGGACATG TGCCATGATA AACTGTCAGT ATAGCTGTGA AGACACAGAA  
GAAGGGCCAC AGTGCCTGTG TCCATCCTCA GGACTCCGCC TGGCCCCAAA  
TGGAAGAGAC TGTCTAGATA TTGATGAATG TGCCTCTGGT AAAGTCATCT  
GTCCCTACAA TCGAAGATGT GTGAACACAT TTGGAAGCTA CTACTGCAAA  
TGTCACATTG GTTTCGAACT GCAATATATC AGTGGACGAT ATGACTGTAT  
AGATATAAAT GAATGTACTA TGGATAGCCA TACGTGCAGC CACCATGCCA  
ATTGCTTCAA TACCCAAGGG TCCTTCAAGT GTAAATGCAA GCAGGGATAT  
AAAGGCAATG GACTTCGGTG TTCTGCTATC CCTGAAAATT CTGTGAAGGA  
AGTCCTCAGA GCACCTGGTA CCATCAAAGA CAGAATCAAG AAGTTGCTTG  
CTCACAAAAA CAGCATGAAA AAGAAGGCAA AAATTAAAAA TGTTACCCCA  
GAACCCACCA GGACTCCTAC CCCTAAGGTG AACTTGCAGC CCTTCAACTA  
TGAAGAGATA GTTTCAGAG GCGGGAAGT TCATGGAGGT AAAAAAGGGA  
ATGAAGAGAA AATGAAAGAG GGGCTTGAGG ATGAGAAAAG AGAAGAGAAA  
GCCCTGAAGA ATGACATAGA GGAGCGAAGC CTGCGAGGAG ATGTGTTTTT  
CCCTAAGGTG AATGAAGCAG GTGAATTCGG CCTGATTCTG GTCCAAAGGA  
AAGCGCTAAC TTCCAAACTG GAACATAAAG ATTTAAATAT CTCGGTTGAC  
TGCAGCTTCA ATCATGGGAT CTGTGACTGG AAACAGGATA GAGAAGATGA  
TTTTGACTGG AATCCTGCTG ATCGAGATAA TGCTATTGGC TTCTATATGG  
CAGTTCCGGC CTTGGCAGGT CACATGAAAG ACATTGGCCG ATTGAAACTT  
CTCCTACCTG ACCTGCAACC CCAAAGCAAC TTCTGTTTGC TCTTTGATTA  
CCGGCTGGCC GGAGACAAAG TCGGGAAACT TCGAGTGTTT GTGAAAAACA  
GTAACAATGC CCTGGCATGG GAGAAGACCA CGAGTGAGGA TGAAAAGTGG  
AAGACAGGGA AAATTCAGTT GTATCAAGGA ACTGATGCTA CAAAAGCAT  
CATTTTGGAA GCAGAACGTG GCAAGGGCAA AACC GGCGAA ATCGCAGTGG  
ATGGCGTCTT GCTTGTTTCA GGCTTATGTC CAGATAGCCT TTTATCTGTG  
GANNNCTGAA TGGTACTATC TTTATATTTG ACTTTGTATG TCAGTTCCCT  
GGTTTTTTTG ATATTGCATC ATAGGACCTC TGGCATTTTA AAATTACTAG  
CTGAAAAATT G  
(SEQ ID NO 2)

FIG. 2

